

# SEQUENCE LISTING

<110> Spencer, David  
Hanks, Brent  
Slawin, Kevin

<120> Induced Activation in Dendritic Cells

<130> P02165US1

<140> Not Assigned

<141> 2004-02-18

<150> US 60/448,046

<151> 2003-02-18

<160> 4

<170> PatentIn version 3.1

<210> 1

<211> 228

<212> DNA

<213> Mouse

<400> 1

tatatcaaaa aggtgggtcaa gaaaccaaag gataatgaga tggtaccccc tgcgggtcga 60

cggcaagatc cccaggagat ggaagattat cccggtcata acaccgctgc tccagtgcag 120

gagacactgc acgggtgtca gcctgtcaca caggaggatg gtaaagagag tcgcatctca 180

gtgcaggagc ggcaggtgac agacagcata gccttgagge ccctggtc 228

<210> 2

<211> 76

<212> PRT

<213> Mouse

<400> 2

Tyr Ile Lys Lys Val Val Lys Lys Pro Lys Asp Asn Glu Met Leu Pro  
1 5 10 15

Pro Ala Ala Arg Arg Gln Asp Pro Gln Glu Met Glu Asp Tyr Pro Gly  
20 25 30

His Asn Thr Ala Ala Pro Val Gln Glu Thr Leu His Gly Cys Gln Pro  
35 40 45

Val Thr Gln Glu Asp Gly Lys Glu Ser Arg Ile Ser Val Gln Glu Arg  
50 55 60

Gln Val Thr Asp Ser Ile Ala Leu Arg Pro Leu Val  
65 70 75

<210> 3  
 <211> 1579  
 <212> DNA  
 <213> Mouse

<400> 3  
 tgccctgcat ggtgtccttg cctcggctgt gcgcgctatg gggctgcttg ttgacagcgg 60  
 tccatctagg gcagtgtgtt acgtgcagtg acaaacagta cctccacgat ggccagtgtc 120  
 gtgatttgtg ccagccagga agccgactga caagccactg cacagctctt gagaagaccc 180  
 aatgccaccc atgtgactca ggcgaattct cagcccagtg gaacagggag attcgctgtc 240  
 accagcacag aactgtgaa cccaatcaag ggcttcgggt taagaaggag ggcaccgcag 300  
 aatcagacac tgtctgtacc tgtaaggaag gacaacactg caccagcaag gattgcgagg 360  
 catgtgtcga gcacacgccc tgtatccctg gctttggagt tatggagatg gccactgaga 420  
 ccactgatac cgtctgtcat ccctgcccag tcggcttctt ctccaatcag tcatcacttt 480  
 tcgaaaagtg ttatccctgg acaagctgtg aggataagaa cttggaggtc ctacagaaag 540  
 gaacgagtca gactaatgtc atctgtgggt taaagtcccg gatgcgagcc ctgctgggtc 600  
 ttctgtcgt gatgggcac ctcacacca ttttcgggggt gtttctctat atcaaaaagg 660  
 tggtaagaa accaaaggat aatgagatgt taccctctgc ggctcgacgg caagatcccc 720  
 aggagatgga agattatccc ggtcataaca ccgctgctcc agtgcaggag aactgcacg 780  
 ggtgtcagcc tgtcacacag gaggatggta aagagagtcg catctcagtg caggagcggc 840  
 aggtgacaga cagcatagcc ttgaggcccc tggctctgaac cctggaactg ctttggaggc 900  
 gatggctgt tgtgtacctt tgaagtttga gatgagccaa gacagagccc agtgcagcta 960  
 actctcatgc ctgccccctg tcatttctca acttgctttt taaggatgga gggaaagctc 1020  
 gggcatcggg aggtccacag tgatatctac caagtgcagc agtgcaggac ccagagttgt 1080  
 cttgtgcgg cgttcactgt aaggagtcgt ggctacagga gtccgtggcc cgcagcttgt 1140  
 gctcgtagag ggcacctggt tgccatcagc agggactgg ctaaataaat ctgtaattat 1200  
 ttatacaatg gcatctcaga aactctagca ggtggggcag aaaacaggta gtggaatgat 1260  
 gggtagagaa acagctttta aaacacattc caaggcagggt aagatggctt ttgtgggtaa 1320  
 aggagcttgc tgcccaaacc cggttacctg attttgatcc ctgggacttc atggtaaaag 1380  
 ggagagaacc aaatccagag ggttgtcatt tgacctccat gtgtgctctg tggtaatgta 1440  
 ccccggtgtg gcacatgtgc acatattcta aaatggatgt ggtgggtgtat ttagaaaatt 1500  
 atttaatccg ccctgggttt ctacctgtgt gttaccattt agttcttgaa taaagacaca 1560  
 ctcaaccttt atatttaca 1579

<210> 4  
 <211> 289  
 <212> PRT  
 <213> Mouse

<400> 4

Met Val Ser Leu Pro Arg Leu Cys Ala Leu Trp Gly Cys Leu Leu Thr  
 1 5 10 15

Ala Val His Leu Gly Gln Cys Val Thr Cys Ser Asp Lys Gln Tyr Leu  
 20 25 30

His Asp Gly Gln Cys Cys Asp Leu Cys Gln Pro Gly Ser Arg Leu Thr  
 35 40 45

Ser His Cys Thr Ala Leu Glu Lys Thr Gln Cys His Pro Cys Asp Ser  
 50 55 60

Gly Glu Phe Ser Ala Gln Trp Asn Arg Glu Ile Arg Cys His Gln His  
 65 70 75 80

Arg His Cys Glu Pro Asn Gln Gly Leu Arg Val Lys Lys Glu Gly Thr  
 85 90 95

Ala Glu Ser Asp Thr Val Cys Thr Cys Lys Glu Gly Gln His Cys Thr  
 100 105 110

Ser Lys Asp Cys Glu Ala Cys Ala Gln His Thr Pro Cys Ile Pro Gly  
 115 120 125

Phe Gly Val Met Glu Met Ala Thr Glu Thr Thr Asp Thr Val Cys His  
 130 135 140

Pro Cys Pro Val Gly Phe Phe Ser Asn Gln Ser Ser Leu Phe Glu Lys  
 145 150 155 160

Cys Tyr Pro Trp Thr Ser Cys Glu Asp Lys Asn Leu Glu Val Leu Gln  
 165 170 175

Lys Gly Thr Ser Gln Thr Asn Val Ile Cys Gly Leu Lys Ser Arg Met  
 180 185 190

Arg Ala Leu Leu Val Ile Pro Val Val Met Gly Ile Leu Ile Thr Ile  
 195 200 205

Phe Gly Val Phe Leu Tyr Ile Lys Lys Val Val Lys Lys Pro Lys Asp

210		215		220
Asn Glu Met Leu Pro Pro Ala Ala Arg Arg Gln Asp Pro Gln Glu Met				
225		230		235 240
Glu Asp Tyr Pro Gly His Asn Thr Ala Ala Pro Val Gln Glu Thr Leu				
	245		250	255
His Gly Cys Gln Pro Val Thr Gln Glu Asp Gly Lys Glu Ser Arg Ile				
	260		265	270
Ser Val Gln Glu Arg Gln Val Thr Asp Ser Ile Ala Leu Arg Pro Leu				
	275		280	285
Val				